

Docket No.: A7542.0000/P001-E (PATENT)

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Patent Application of:

Ginette Serrero

Application No.: Not Yet Assigned

Group Art Unit: 1642

Filed: April 4, 2001

Examiner: M. Wells

For: 88KDA TUMORIGENIC GROWTH

FACTOR AND ANTAGONISTS

REQUEST TO USE COMPUTER READABLE FORM FROM ANOTHER APPLICATION

ATTN: Application Processing Division Assistant Commissioner for Patents

Washington, DC 20231

Dear Sir:

The computer readable form in this divisional application, filed April 4, 2001, is identical with that filed in application numbers 08/991,862 filed on December 16, 1997 which is a continuation-in-part application of U.S. application serial number 08/863,079 filed on May 23, 1997. In accordance with 37 CFR 1.821(e), please use the only computer readable form filed in the applications as the computer readable form for the instant application. It is understood that the Patent and Trademark Office will make the necessary change in application number and filing date for the computer readable form that will be used for the instant application. A paper copy of the Sequence Listing is included in a separately filed preliminary amendment for incorporation into the specification.

Dated: April 4, 2001

Respectfully submitted,

James W. Brady, Jr

Registration No.: 32,115

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Registration No.: 40,399

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Attorneys for Applicant



Atty. Docket No: Z9996.488/PQ01-A

In re patent application of
Serrero, Ginette

Serial No. 08/991,862

Filed: December 16, 1997

For: 88 KDA TUMORIGENIC GROWTH FACTOR AND ANTAGONISTS

STATEMENT TO SUPPORT FILING AND SUBMISSION IN ACCORDANCE WITH 37 C.F.R. §§ 1.821-1.825

Assistant Commissioner for Patents Washington, D.C. 20231
Box SEQUENCE

Sir:

In connection with a Sequence Listing submitted concurrently herewith, the undersigned hereby states that:

- the submission, filed herewith in accordance with 37
 C.F.R. § 1.821(g), does not include new matter;
- 2. the content of the attached paper copy and the attached computer readable copy of the Sequence Listing, submitted in accordance with 37 C.F.R. § 1.821(c) and (e), respectively, are the same; and
- 3. all statements made herein of their own knowledge are true and that all statements made on information and belief are believed to be true; and further, that these statements were made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United

.

States Code and that such willful false statements may jeopardize the validity of the application or any patent resulting therefrom.

Respectfully submitted,

Date

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Intellectual Property Services
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Portsmouth, N.H.
800-318-3021

2



SEQUENCE LISTING



<110> Serrero, Ginette

<120> 88 KDA TUMORIGENIC GROWTH FACTOR AND ANTAGONISTS

<130> Z9996.488/P001-A

<140> 08/991,862

<141> 1997-12-16

<150> 08/863,862

<151> 1997-05-23

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<170> PatentIn Ver. 2.0

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<213> Mouse epithelin/granulin

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<222> (23)..(1789)

<223> The sequence is identical to that of the published mouse granulin except for one nucleotide (T instead of G) at position 1071 of GP88 cDNA (position 1056 of mouse granulin).

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cct gtt gcc tgc tgc ctt gac cag gga gga gcc aac tac agc tgc tgt 148 Pro Val Ala Cys Cys Leu Asp Gln Gly Gly Ala Asn Tyr Ser Cys Cys 30 30 35

aac cct ctt ctg gac aca tgg cct aga ata acg agc cat cat cta gat 196
Asn Pro Leu Leu Asp Thr Trp Pro Arg Ile Thr Ser His His Leu Asp
45 50 55

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Gly Ser Cys Gln Thr His Gly His Cys Pro Ala Gly Tyr Ser Cys Leu
60 65 70

ctc act gtg tct ggg act tcc agc tgc tgc ccg ttc tct aag ggt gtg
Leu Thr Val Ser Gly Thr Ser Ser Cys Cys Pro Phe Ser Lys Gly Val
75 80 85 90

tct tgt ggt gat ggc tac cac tgc tgc ccc cag ggc ttc cac tgt agt
Ser Cys Gly Asp Gly Tyr His Cys Cys Pro Gln Gly Phe His Cys Ser
95 100 105

·" .								. •		•		•			
gca ga Ala As	t gg p Gl	g aa y Ly 11	s Ser	cgc Cys	ttc Phe	Gln	atg 1 Met : 115	tca Ser	gat Asp	aac Asn	ccc Pro	Leu 120	ggt Gly	gct Ala	388
gtc ca Val Gl	g to n Cy 12	s Pr	t ggg o Gly	g agc / Ser	cag Gln	ttt Phe 130	gaa Glu	tgt Cys	cct Pro	gac Asp	tct Ser 135	gcc Ala	acc Thr	tgc Cys	436
tgc at Cys Il	e Me	g gt et Va	t gan 1 Ası	ggt Gly	tcg Ser 145	tgg Trp	gga Gly	tgt Cys	tgt Cys	ccc Pro 150	atg Met	ccc Pro	cag Gln	gcc Ala	484
tct to Ser Cy 155	gc to ys C	gt ga ys Gl	ia ga .u As	c aga p Arg 160	Val	cat His	tgc Cys	tgt Cys	ccc Pro 165	cat His	ggg Gly	gcc Ala	tcc Ser	tgt Cys 170	532
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aat g Asn A 235	cc a la l	atc t [le C	gc to ys C	gt tc ys Se 24	r Asp	c cac	ctg Leu	cac	Cys 245	Cys	ccc Pro	c cag	gac Asp	act Thr 250	772
gta t Val (gt q Cys A	gac c Asp I	eu I	tc ca le Gl 55	g agt n Sei	t aaq r Lys	g tgc s Cys	cta Lev 260	ı Şei	c aaq r Lys	g aad s Asi	tac n Tyr	acc Thr 265	rnr	820
gat (Asp 1	ctc (Leu :	Leu 1	acc a Thr L 270	ag ct ys Le	g cc u Pr	t gga o Gl	a tac y Tyr 275	Pro	a gto o Vai	g aad l Ly:	g gad s Gl	g gto u Val 280	LLy	g tgc s Cys	868
gac a Asp I	Met	gag (Glu 1 285	gtg a Val S	gc to er Cy	jc cc 's Pr	t ga o Gl 29	u Gly	a tai	t ac	c tg r Cy	c tg s.Cy 29	s Ar	c cte	c aac u Asn	916
Thr	ggg Gly 300.	gcc Ala	tgg g Trp G	gc to	gc tg ys Cy 30	s Pr	a tti o Phe	t gc	c aa a Ly	g gc s Al 31	a Va	gʻtg 1 Cy	t tg s Cy	t gac s Asp	964
gat Asp 315	cac His	att Ile	cat t His (Cys C	gc cc ys Pr 20	g go	a gg a Gl	g tt y Ph	t ca e Gl 32	.n Cy	t ca 's Hi	c ac s Th	a ga r Gl	gʻaaa u Lys 330	1012
gga Gly	acc Thr	tgc Cys	Glu I	atg g Met G 335	gt at ly I]	ic ct le Le	c ca eu Gl	a gt n Va 34	1 G1	gg tg Ly Tr	gg at p Me	g aa et Ly	ig aa 's Ly 34	ig gtc vs Val 15	1060

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agg Arg	aco Thi	tgi Cy:	t gaq s Glu	g aac 1 Lys 495	Asp	gtc Val	gat Asp	ttt Phe	atc Ile 500	e Gli	g cct n Pro	t cco	c gte o Va	g ct l Le 50	c ctg u Leu 5	1540
acc Thi	c cto	c gg u Gl	c cct y Pro	o Lys	g gtt s Val	ggç Gl	g aat 7 Asr	gtg Val	. Gl	g tg	t gg s Gl	a ga y Gl	a gg u Gl 52	у Ні	t ttc s Phe	1588
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tg: Cy:	c tg s Cy 54	s Pr	c ta o Ty	c ct r Le	a aaq u Ly:	g gg s Gl ₂ 54	y Va	c tgo l Cys	c tg s Cy	t ag s Ar	a ga g As 55	p G1	a cg y Ar	gt ca ng Hi	ac tgt is Cys	1684
tg Cy 55	s Pr	c gg	jt gg .y Gl	c tt y Ph	c cade Hi	s Cy	t tc s Se	a gce r Al	c ag a Ar	g gg g Gl 56	y Th	c aa nr Ly	ng to	gt ti /s Le	cg• cga eu Arg 570	1732
aa Ly	g aa s Ly	ig at 's II	t co le Pr	t cg o Ar 57	g Tr	g ga p As	c at p Me	g tt t Ph	t tt e Le 58	u Ar	gg ga cg As	at co sp Pi	og gt ro Va	al P	ca aga ro Arg 85	1780

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Trp Pro Arg Ile Thr Ser His His Leu Asp Gly Ser Cys Gln Thr His 50 55 60

Gly His Cys Pro Ala Gly Tyr Ser Cys Leu Leu Thr Val Ser Gly Thr 65 70 75 80

Ser Ser Cys Cys Pro Phe Ser Lys Gly Val Ser Cys Gly Asp Gly Tyr 85 90 95

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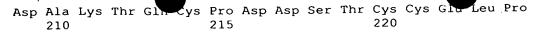
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Val His Cys Cys Pro His Gly Ala Ser Cys Asp Leu Val His Thr Arg 165 170 175

Cys Val Ser Pro Thr Gly Thr His Thr Leu Leu Lys Lys Phe Pro Ala 180 185 190

Gln Lys Thr Asn Ser Ala Val Ser Leu Pro Phe Ser Val Val Cys Pro 195 200 205



Thr Gly Lys Tyr Gly Cys Cys Pro Met Pro Asn Ala Ile Cys Cys Ser 225 230 235 240

Asp His Leu His Cys Cys Pro Gln Asp Thr Val Cys Asp Leu Ile Gln 245 250 255

Ser Lys Cys Leu Ser Lys Asn Tyr Thr Thr Asp Leu Leu Thr Lys Leu 260 265 270

Pro Gly Tyr Pro Val Lys Glu Val Lys Cys Asp Met Glu Val Ser Cys 275 280 285

Pro Glu Gly Tyr Thr Cys Cys Arg Leu Asn Thr Gly Ala Trp Gly Cys 290 295 300

Cys Pro Phe Ala Lys Ala Val Cys Cys Asp Asp His Ile His Cys Cys 305 310 315

Pro Ala Gly Phe Gln Cys His Thr Glu Lys Gly Thr Cys Glu Met Gly 325 330 335

Ile Leu Gln Val Gly Trp Met Lys Lys Val Ile Ala Pro Leu Arg Leu 340 345 350

Pro Asp Pro Gln Ile Leu Lys Ser Asp Thr Pro Cys Asp Asp Phe Thr 355 360 365

Arg Cys Pro Thr Asn Asn Thr Cys Cys Lys Leu Asn Ser Gly Asp Trp 370 375. 380

Gly Cys Cys Pro Ile Pro Glu Ala Val Cys Cys Ser Asp Asn Gln His

Cys Cys Pro Gln Gly Phe Thr Cys Leu Ala Gln Gly Tyr Cys Gln Lys 405 410 415

Gly Asp Thr Met Val Ala Gly Leu Glu Lys Ile Pro Ala Arg Gln Thr 420 425 430

Thr Pro Leu Gln Ile Gly Asp Ile Gly Cys Asp Gln His Thr Ser Cys 435 440 445

Pro Val Gly Gln Thr Cys Cys Pro Ser Leu Lys Gly Ser Trp Ala Cys 450 455 460

Cys Gln Leu Pro His Ala Val Cys Cys Glu Asp Arg Gln His Cys Cys 465 470 475 480

Pro Ala Gly Tyr Thr Cys Asn Val Lys Ala Arg Thr Cys Glu Lys Asp 485 490 495

Val Asp Phe Ile Gln Pro Pro Val Leu Leu Thr Leu Gly Pro Lys Val 500 505 510

Gly Asn Val Glu Cys Gly Glu Gly His Phe Cys His Asp Asn Gln Thr 515 520 525

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                       535
Gly Val Cys Cys Arg Asp Gly Arg His Cys Cys Pro Gly Gly Phe His
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                    550
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 Ser Asp Thr
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                    5
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  <221> PEPTIDE
  <222> (1)..(14)
  <223> Internal peptide of mouse GP88 used to raise the
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Same and the second

antisera against the GP88 used in the

immunoaffinity step.

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                   5
                                      10
Arg Asp Val
 <210> 7
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  <221> primer
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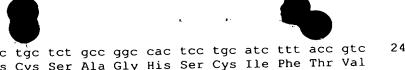
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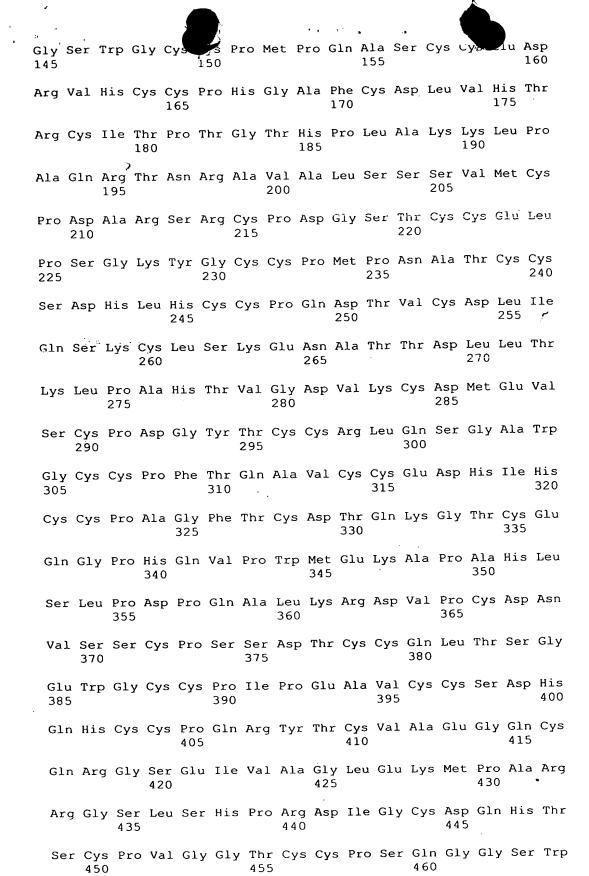
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tgc tgc ctg gac ccc gga gga gcc agc tac agc tgc tgc cgt ccc ctt Cys Cys Leu Asp Pro Gly Gly Ala Ser Tyr Ser Cys Cys Arg Pro Leu 30 35 40	147
ctg gac aaa tgg ccc aca aca ctg agc agg cat ctg ggt ggc ccc tgc Leu Asp Lys Trp Pro Thr Thr Leu Ser Arg His Leu Gly Gly Pro Cys 50 55 60	: 195



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tca Ser	ggg Gly	act Thr 80	tcc Ser	agt Ser	tgc Cys	tgc Cys	ccc Pro 85	ttc Phe	cca Pro	gag Glu	gcc Ala	gtg Val 90	gca Ala	tgc Cys	G.	J y	291	
gat Asp	ggc Gly 95	cat His	cac His	tgc Cys	tgc Cys	cca Pro 100	cgg Arg	ggc Gly	ttc Phe	cac His	tgc Cys 105	agt Ser	gca Ala	gac Asp	g G	gg ly	339	
cga Arg 110	tcc Ser	tgc Cys	ttc Phe	caa Gln	aga Arg 115	tca Ser	ggt Gly	aac Asn	aac Asn	tcc Ser 120	gtg Val	ggt Gly	gcc Ala	ato Ile	3 6	ag ln 25	387	
tgc Cys	cct Pro	gat Asp	agt Ser	cag Gln 130	ttc Phe	gaa Glu	tgc Cys	ccg Pro	gac Asp 135	Phe	tcc Ser	acg Thr	tgc Cys	tg: Cy: 14	5 V	tt al	435	
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gt. Va	c ato l Me	g tg t Cy	t cc s Pr	g gad o As _l 21	c gca p Ala 0	a cgg	g tco g Sei	c cgo	g tg g Cy 21	s Pr	t ga o As	t gg p Gl	t tc y Se	r T	ec nr 20	tgc Cys	675	
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cac His	ata Ile	cac His 320	tgc Cys	tgt Cys	ccc Pro	Ala	ggg Gly 325	ttt Phe	acg Thr	tgt Cys	gac Asp	acg Thr 330	cag Gln	aag Lys	ggt Gly	1011
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acg Thr	tct Ser	Gly	gag Glu 385	tgg Trp	ggc Gly	tgc Cys	tgt Cys	cca Pro 390	atc Ile	cca Pro	gag Glu	gct Ala	gtc Val 395	tgc Cys	tgc Cys	1203
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cct Pro 430	Ala	cgc Arg	cgc g Arg	ggt Gly	tcc Ser 435	Leu	tcc Ser	cac His	ccc Pro	aga Arg 440	Asp	ato	ggc Gly	tgt Cys	gac Asp 445	1347
caç Glr	g cad n His	c acc	c ago c Sei	c tgc c Cys 450	Pro	g gtg Val	ggc Gly	gga Gly	acc Thr 455	Cys	tgo Cys	c cco	g ago Sei	c caq Gli 460	g ggt n Gly O	1395
ggo Gl	g ago y Se:	c tgo	g gco p Ala 46	a Cys	c tgo s Cys	c caç s Glr	g ttg n Lei	g ccc i Pro 470	His	gct Ala	gto a Val	g tgo l Cy:	c tgo s Cy: 47	s GI	g gat u Asp	1443
cg Ar	c ca g Gl:	g ca n Hi 48	s Cy	c tgo s Cy:	c cco	g gct o Ala	gg0 a Gl;	у Туі	acc Thi	c tgo	c aad s Asi	c gto n Va 49	l Ly	g gc s Al	t cga a Arg	1491
tc Se	c tg r Cy 49	s Gl	g aa u Ly	g ga s Gl	a gte u Va	g gto 1 Va. 50	l Se	t gcd r Ala	c cad a Gli	g cci n Pro	t gc o Al 50	a Th	c tt r Ph	c ct e Le	g gcc u Ala	1539
cg Ar 51	g Se	c cc r Pr	t ca o Hi	c gt s Va	g gg 1 Gl 51	y Va	g aa l Ly	g gad s As	c gto p Va	g ga 1 G1 52	u Cy	t gg s Gl	g ga y Gl	a gg u Gl	a•cac y His 525	1587
t t Ph	c tg e Cy	c ca s Hi	it ga .s As	it aa sp As 53	n Gl	g ac n Th	c tg r Cy	c tg s Cy	c cg s Ar 53	g As	c aa p As	ic cg in Ar	a ca g Gl	n Gl	gc tgg Ly Trp 10	1635

gcc to									•								
Ala C	gc t ys C	ys F	cc tro 1	ac g Tyr A	ec c	ag g Sln G	Sly V	tc t al C 50	gt t ys C	gt o ys <i>F</i>	gct q Ala <i>H</i>	Asp .	cgg Arg 555	Àrg	са Ні	.s	1683
tgc t Cys C	ys P	ct c ro F	jct (Mla (ggc t Gly 1	tc c Phe <i>F</i>	Arg (gc g Cys A	jca c	gc a rg A	gg (Sly	acc Thr 570	aag Lys	tgt Cys	tt Le	:g eu	1731
cgc a Arg A 5	gg g rg G 75	ag q Slu <i>H</i>	gcc Ala	ccg (Pro i	Arg '	tgg (Trp /	gac g Asp <i>P</i>	gcc o Ala E	ct t ro L	eu i	agg Arg 585	gac Asp	cca Pro	gcc	tt Le	ig eu	1779
aga c Arg G 590				tgag	ggac	ag t	actga	aagao	tct	gca	gccc	tcg	gga	cccc	=		1831
actcg	gago	gg t	gccc	tctg	c tc	aggc	ctcc	cta	gcaco	cțc	cccc	taac	ca	aatt	ct	ccct	1891
ggaco	ccat	tt c	tgag	ctcc	с са	tcac	catg	gga	ggtg	3 99	cctc	aato	ta	aggo	ccc	ttcc	1951
ctgt	cagai	ag g	gggt	tgag	g ca	aaag	ccca	tta	caag	ctg	ccat	ccc	ctc	ccc	gtt	tcag	2011
tggad	cct	gt g	gcca	ıggtg	c tt	ttcc	ctat	cca	cagg	ġgt	gttt	gtg	gt	tgg	gtg	tgct	2071
ttcaa	ataa	ag t	ttgt	cact	t to	:tt										٠	2095
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<212 <213			GP8	B cDi	ΑI												
	> Hu > 17	man				Trp	Val	Ala	Leu 10	Thr	Ala	Gly	Lei	Va 1	ıl <i>I</i>	Ala	
<213 <400 Met	> Hu > 17 Trp	man Thr	Leu	Val 5 Pro	Ser				10					s C)	.5		
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Ala Cys Cys Gln Leu G His Ala Val Cys Cys Glu Asp 465

Gus Cys Bro Ala Cly Tyr Thr Cys Asp Val Lys Ala Arg Ser Cys Glu

Cys Cys Pro Ala Gly Tyr Thr Cys Asn Val Lys Ala Arg Ser Cys Glu 485 490 495

Lys Glu Val Val Ser Ala Gln Pro Ala Thr Phe Leu Ala Arg Ser Pro 500 505 510

His Val Gly Val Lys Asp Val Glu Cys Gly Glu Gly His Phe Cys His 515 520 525

Asp Asn Gln Thr Cys Cys Arg Asp Asn Arg Gln Gly Trp Ala Cys Cys 530 535 540

Pro Tyr Ala Gln Gly Val Cys Cys Ala Asp Arg Arg His Cys Cys Pro 545 550 555 560

Ala Gly Phe Arg Cys Ala Arg Arg Gly Thr Lys Cys Leu Arg Arg Glu
565 570 575

Ala Pro Arg Trp Asp Ala Pro Leu Arg Asp Pro Ala Leu Arg Gln Leu 580 585 590

Leu